# Synchrotron Radiation in Biotechnology:

Combining Physics, Biology, Chemistry, and Mathematics to do Science

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> > Brookhaven Biology

### The Plan for this Lecture

I'm going to take you through some of the adventures and successes scientists in the Brookhaven Biology Department have had in using Brookhaven's National Synchrotron Light Source to make discoveries that have medical or agricultural value.

For example....

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# Lyme disease is an increasing health concern on Long Island.

It affects more than 10,000 Americans each year, and it can be difficult to diagnose. Since it can cause severe complications if untreated, there is a real need for a safe and effective vaccine. This image shows the structure of an

important protein that's located on the outer surface of the bacterium that causes Lyme disease. It's called *OspA*.



To learn the structure of OspA, researchers in **Brookhaven's Biology Department** studied protein *crystals* like these using *x-ray diffraction*.

OspA was used in firstgeneration **Lyme disease vaccines** -- but differences among strains of bacteria can make a vaccination fail.

New models of OspA's structure show exactly where different strains of bacteria vary.

Researchers can use this information to develop secondgeneration vaccines that will give better protection again Lyme disease.







This study of the Lyme Disease
Vaccine is just one example of the way
Brookhaven Biologists are using
Biology, Chemistry, Physics, and Math
to answer questions that can be important to
human health, to agriculture, and to
fundamental science.

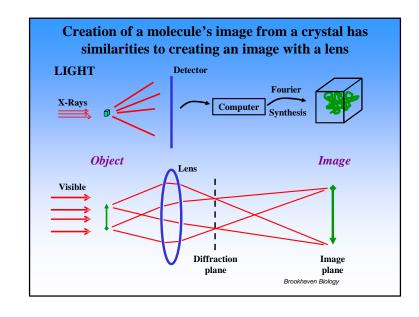
Let's look at how all these disciplines can fit together to provide this wealth of information....

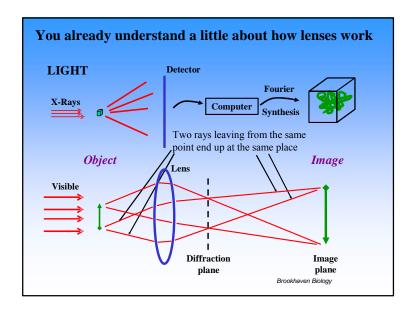
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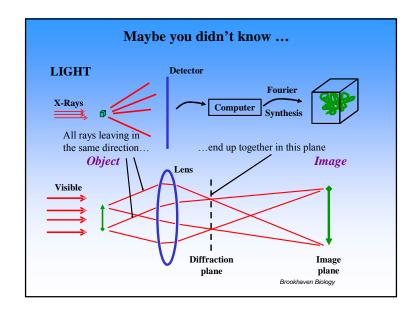
### How does it work?

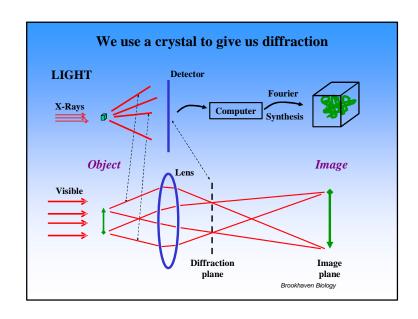
I've told you that we use crystals and x-rays. These allow us to "look" at large molecules like OspA.

You'll be surprised how much you already know about how we might do this.









# Why do we use x-rays?

- The features we're trying to see are on the order of the distance between atoms: 10<sup>-10</sup> meters.
- To "see" the atoms, we need to use light with a wavelength that is near to this distance.
- X-Rays (x-ray light) have a suitable wavelength.

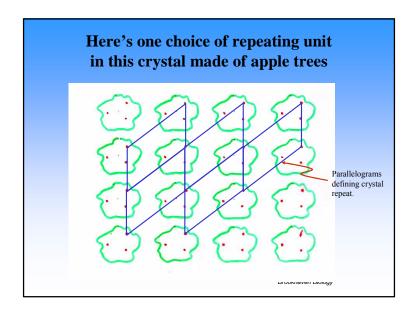
# What is a crystal?

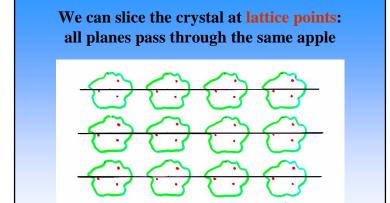
- A crystal is a periodic arrangement of objects (molecules) repeating in two or three dimensions.
- The repeating unit is a parallelepiped (three-dimensional) or a parallelegram (two-dimensional).
- A crystal like that of OspA will be half a mm on a side and contain 10<sup>15</sup> molecules.

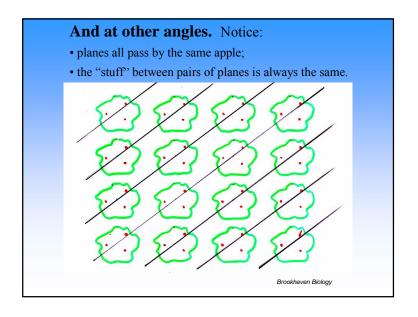
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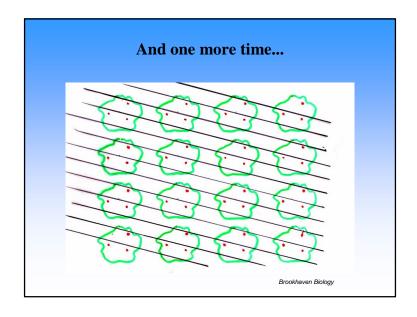
# Why do we use crystals when we'd like to see one molecule?

- We can't focus enough x-rays into a small enough volume to "see" a molecule. We use lots of molecules in a crystal to get a bigger target.
- Even if we could, the x-rays would burn up the molecule.
- Even if that would work, we don't have a lens for the x-rays.
- The crystal amplifies the signal, and gives us a way to get the phase information back.

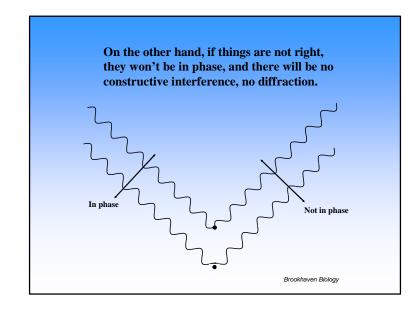


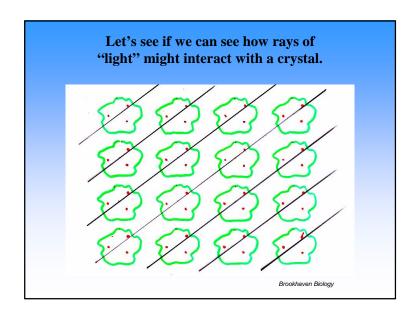


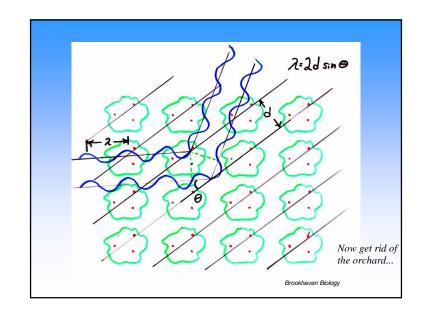


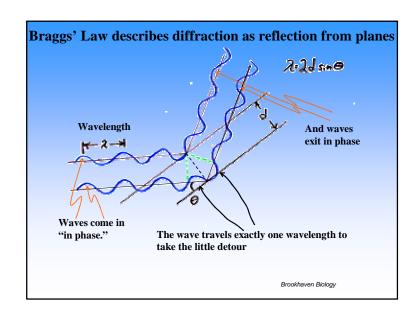


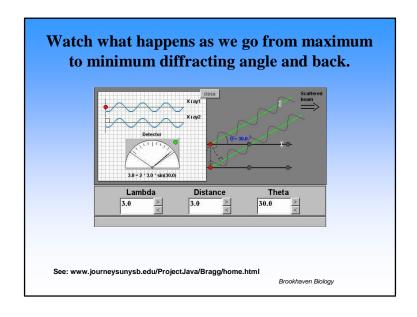
# Diffraction Let's do a thought experiment. Send a beam of x-rays at a pair of single atoms, suspended in space. If the angle is just right for the wavelength and distance between the atoms, the scattered x-rays will be in phase, and they will interfere constructively. In phase Brookhaven Biology

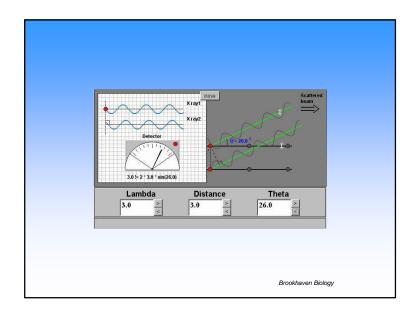


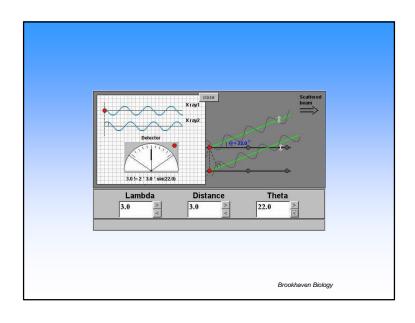


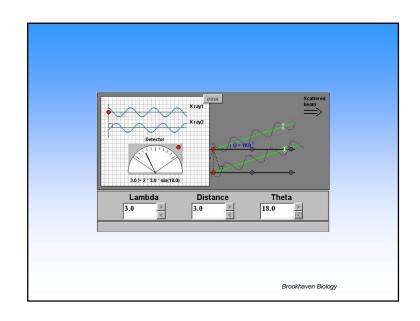


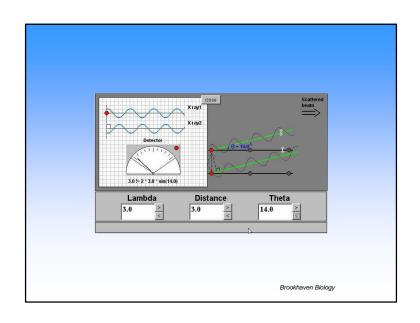


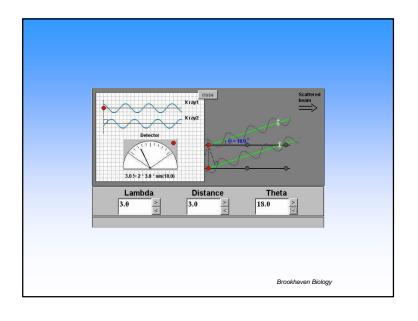


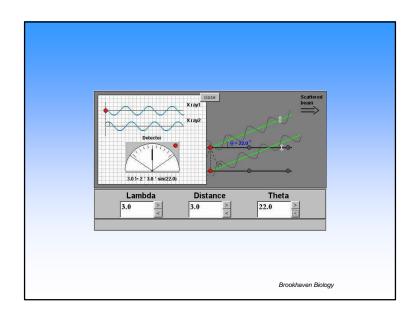


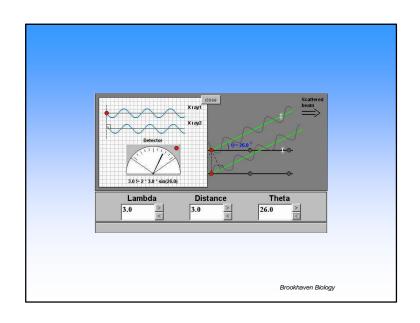


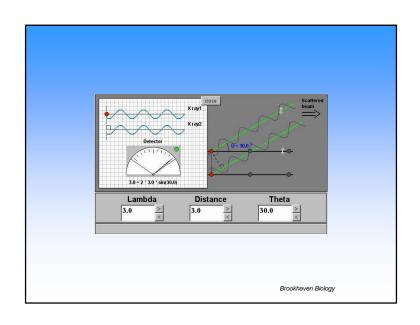


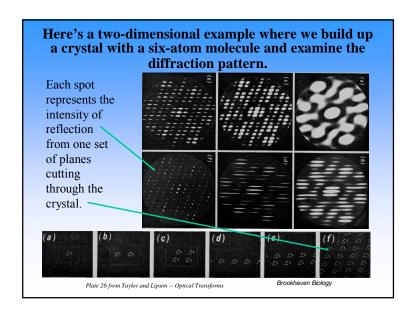


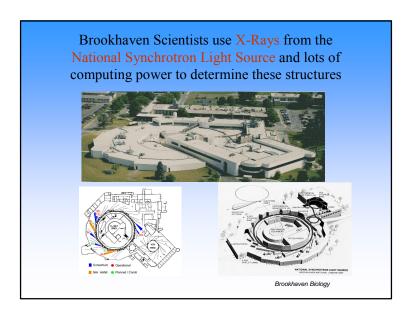


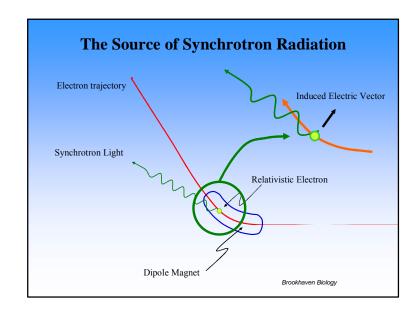


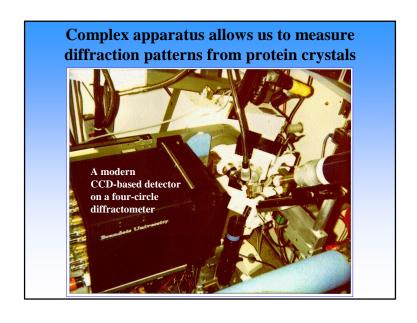


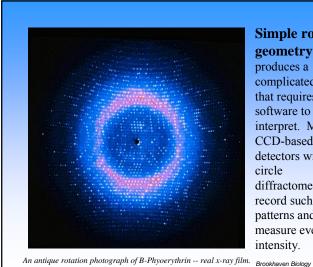












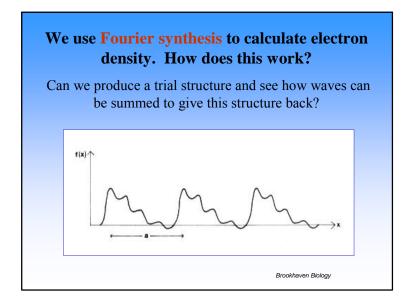
**Simple rotation** geometry produces a complicated pattern that requires good software to interpret. Modern CCD-based detectors with fourcircle diffractometers record such patterns and measure every spot intensity.

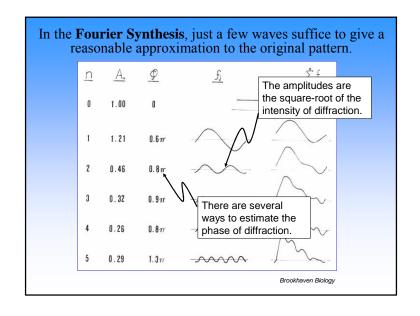
**Q**: How do we perform the second interference step in the functioning of the lens -- to reconstruct the image of the original object?

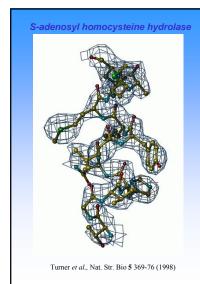
A: We will have to calculate it.

**Q**: How will we represent that object? A: The x-rays are scattered from electrons in the atoms of the crystal.

**Therefore**: for us, the "image" is going to be a representation of the electron density.







Typically, we represent regions of high electron density with "chickewire" cages like this.

The molecular model can be built to fit inside.

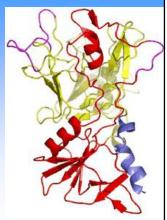
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# Here are a couple more examples...

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## Malaria is a serious world-wide health problem.

It infects approximately 400 million people worldwide, and kills about two million people each year — mostly children. This image shows the structure of an important protein that's located on the outer surface of the parasite that causes malaria. It's called *Apical Membrane Antigen-1*.

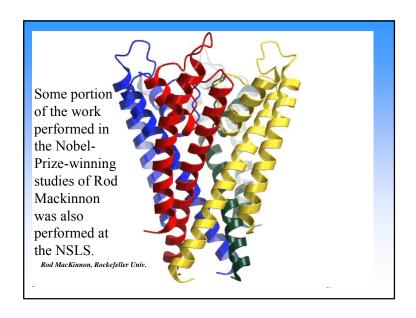


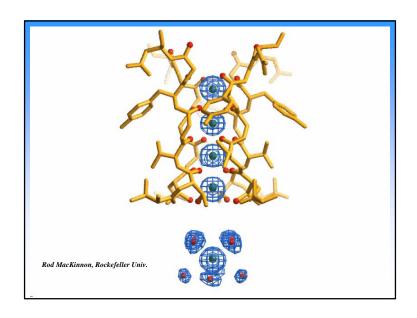
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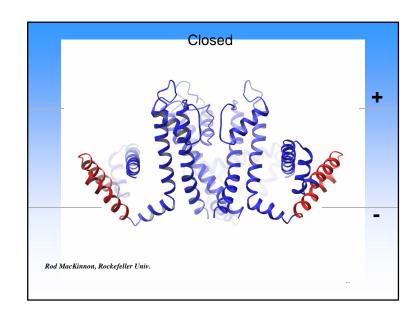
# Strategies for treatment of malaria

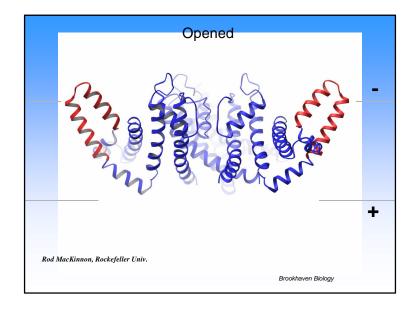
Brookhaven's Mike Becker is part of the team that solved this structure. This molecule itself may be an effective part of a vaccine. In addition, features of the molecule suggest that it might be target for a drug that might be designed to bind to it.











# **Final Thoughts:**

- We are in a golden age for biomedical research.
- Structural biology plays an important role by providing a tangible image of life processes.
- No single technique provides all the answers; we must collaborate.
- Synchrotron x-ray sources are a crucial component.
- We also depend on collaboration among National funding agencies.





